

Table 7. Primers

5	SEQ ID NO. 96	GPT	5'-GTTTAACCACTGCATGATG-3'
	SEQ ID NO. 97	TK _R	5'-GTCCCATCGAGTGGGTAC-3'
	SEQ ID NO. 98	GLY1	5'-CGTGACATGGTACATTCGGACACTTGGCGCACTTCATAAGCGGA-3'
	SEQ ID NO. 99	GLY2	5'-TGCCTCATACACAAATGGAGCTCTGGGACGAGTCGTTCGTGAC-3'
	SEQ ID NO. 100	GLY3	5'-TACCCAGCAGCGGGAGCTCTGTTCCTCCGAACGACGGGCAC-3'
	SEQ ID NO. 101	GLY4	5'-TGTCGTGGTGGGACGGAGGCCCTGCCCTAGCTGCGAGCGTGGG-3'
10	SEQ ID NO. 102	GLY5	5'-CGTTATGTGGCCCGGTAGATTGAGCACTGGCAGTCTGCACCGTCTC-3'
	SEQ ID NO. 103	GLY6	5'-CAGGGCCGTTGTAGGCCCTCCCACTGCATCATATCCCAAGC-3'
	SEQ ID NO. 104	OVR1	5'-CCGGAATGTACCATGTACGAACGAC-3'
	SEQ ID NO. 105	OVR2	5'-GCCTCATTGTGTATGAGGACGCGG-3'
	SEQ ID NO. 106	OVR3	5'-GAGCTCCGCTGCTGGGTAGCGC-3'
	SEQ ID NO. 107	OVR4	5'-CCCTCCGTCCCAACCAACGACAATACG-3'
15	SEQ ID NO. 108	OVR5	5'-CTACCCGGGGCCACATAACGGGTCACCG-3'
	SEQ ID NO. 109	OVR6	5'-GGAGGCCCTACAAACGGCCCTGGTGG-3'
	SEQ ID NO. 110	GPT-2	5'-TTCTATCGATTAAATAGAATTC-3'
	SEQ ID NO. 111	TK _R -2	5'-GCCATACGCTCACAGCCGATCCC-3'

nucleotides underlined represent additional restriction site

nucleotides in bold represent mutations with respect to the original HCC10A sequence

original
paper
copy,
highlighted
letters.
bold

1FW
copy

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SEQ ID NO. 96	GPT	5'-GTTTAACTGCTGATG-3'
SEQ ID NO. 97	TK _n	5'-GTCCCATCGAGTGGGCTAC-3'
SEQ ID NO. 98	GLY1	5'-CGTGACATGGTACATCCGGACACTTGGCGCACTTCATAAGCGGA-3'
SEQ ID NO. 99	GLY2	5'-TGCCCTCATACACAATGGAGCTCTGGGACGAGTCGTTCTGTGAC-3'
SEQ ID NO. 100	GLY3	5'-TACCCAGCAGCGGGAGCTCTGTTGCTCCCGAACGCAGGGGCAC-3'
SEQ ID NO. 101	GLY4	5'-TGTCGTGGTGGGACGGAGGCTTGCCTAGCTGCGAGCGTGGG-3'
SEQ ID NO. 102	GLY5	5'-CGTTATGTGGCCCGGTTAGATTGAGCAGTGGCAGTCCCTGCACCGTCTC-3'
SEQ ID NO. 103	GLY6	5'-CAGGGCCGTTGTAGGCCCTCCACTGCATCATCATATCCCAAGC-3'
SEQ ID NO. 104	OVR1	5'-CCGGAAATGTACCATGTACGAACGAC-3'
SEQ ID NO. 105	OVR2	5'-GCTCCATTGTGTATGAGGACGCGG-3'
SEQ ID NO. 106	OVR3	5'-GAGCTCCCGCTGCTGGGTAGCGC-3'
SEQ ID NO. 107	OVR4	5'-CCCTCCGTCCCCACCCACGACAAATACG-3'
SEQ ID NO. 108	OVR5	5'-CTACCCGGGGCCACATAACGGGTCACCCG-3'
SEQ ID NO. 109	OVR6	5'-GGAGGCCCTACAAACGGCCCTGGTGG-3'
SEQ ID NO. 110	GPT-2	5'-TTCTATCGATTAAATAGAATTC-3'
SEQ ID NO. 111	TK _n -2	5'-GCCATACGCTCACAGCCGATCCC-3'

nucleotides underlined represent additional restriction site

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